Sequence Seq

US-10-100-957A-170 US-10-072-036-113 US-10-072-036-115 US-10-072-036-159 US-10-221-461-11 US-10-221-461-11 US-10-072-036-41 US-10-072-036-41 US-10-072-036-42 US-10-072-036-43 US-10-072-036-43 US-10-072-036-45 US-10-072-036-45 US-10-072-036-45 US-10-072-036-45 US-09-554-000-6 US-09-554-000-6 US-09-554-000-6 US-10-221-461-14 US-10-072-036-125

US-10-072-036-71

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June 3, 2003, 15:08:45; search time 45 Seconds (without alignments) 537.610 Million cell updates/sec
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1274
1 WVSKGEELFTGVVPILVELD......VLLGFVTAAGITLGMDELYR 239
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/ cgn2_6/ptodatta/1/pubpaa/PCT_WEW_PUB.pep:*
/ cgn2_6/ptodatta/1/pubpaa/NSO6_NEW_PUB.pep:*
/ cgn2_6/ptodatta/1/pubpaa/USO6_NEW_PUB.pep:*
/ cgn2_6/ptodatta/1/pubpaa/USO7_NEW_PUB.pep:*
/ cgn2_6/ptodatta/1/pubpaa/USO7_PUBCOMB.pep:*
/ cgn2_6/ptodatta/1/pubpaa/USO7_PUBCOMB.pep:*
/ cgn2_6/ptodatta/1/pubpaa/USO9_NEW_PUB.pep:*
/ cgn2_6/ptodatta/1/pubpaa/USO9_NEW_PUB.pep:*
/ cgn2_6/ptodatta/1/pubpaa/USO9_NEW_PUB.pep:*
GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
                                                                                                         - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                       BLOSUM62
Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 2000000000
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Perfect score:
Sequence:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

/cgn2_6/ptodata/l/pubpaa/US10_NEW_PUB.pep:* /cgn2_6/ptodata/l/pubpaa/US10_PUBCOMB.pep:* /cgn2_6/ptodata/l/pubpaa/US60_NEW_PUB.pep:* /cgn2_6/ptodata/l/pubpaa/US60_PUBCOMB.pep:*

Description		Sequence 4, Appli	Sequence 2, Appli	Sequence 4, Appli	4, 4	Sequence 2, Appli	Sequence 13, Appl	Sequence 7, Appli	Sequence 46, Appl	7	٧,	35,	33,	Sequence 34, Appl	Sequence 129, App	Sequence 32, Appl	Sequence 30, Appl	Sequence 31, Appl	Sequence 29, Appl	Sequence 127, App
	Desc	Sequ	Sedi	Sedi	Sedi	Sedi	Sedi	Sedi	Sed	Se.	Seq	Seq	Sedi	Sedi	Sed	Sedi	Sedi	Sedi	Sedi	Sedi
	٠	:																		
	ID	US-09-887-784-4	US-09-887-784-2	US-09-999-745-4	US-09-866-538-4	US-09-797-496B-2	US-10-121-258-13	US-10-221-461-7	US-10-100-957A-46	US-09-920-922-2	US-10-100-957A-2	US-10-033-717-35	US-10-033-717-33	US-10-033-717-34	US-10-072-036-129	US-10-033-717-32	US-10-033-717-30	US-10-033-717-31	US-10-033-717-29	US-10-072-036-127
	DB	6	σ	6	6	σ	Φ	6	σ	10	6	σ	Φ	6	6	σ	σ	Ø	σ	σ
	Query Match Length DB	239	239	239	239	239	239	239	239	239	294	308	359	359	379	391	397	403	429	442
æ	Query Match	100.0	99.4	1.66	99.1	99.1	99.1	99.1	99.1	1.66	99.1	99.1	1.66	99.1	99.1	99.1	1.66	99.1	99.1	99.1
	Score	1274	1266	1263	1263	1263	1263	1263	1263	1263	1263	1263	1263	1263	1263	1263	1263	1263	1263	1263
	Result No.	П	7	m	4	'n	9	7	80	6	10	11	12	13	7.	15	16	17	18	19

RESULT 2 US-09-887-784-2 ; Sequence 2, Application US/09887784

ength 239; ndels 0; Gaps 0;	JILKFICTTGKLPVPWPT 60 	DGNYKTRAEVKFEGDTL 120 	(VNFKIRHNIEDGSVQLA 180 	SFVTAAGITLGMDELYK 239 	
Score 1274; DB 9; Pred. No. 4.8e-115;); Mismatches 0;	ELDGDVNGHKFSVSGEGEGDATYGKI 	HMKQHDFFKSAMPEGYVOERTIFFKU 	LGHKLEYNYNSHNYY IMADKQKNG IP 	'DNHYLSTQSALSKDPNEKRDHWVLLG 	
100.0 100.0	VPILA VPILA	SRYPI SRYPI	EDGN	PVLLI	
20	LFTGV LFTGV	GVQCE GVQCE	GIDFR GIDFR	PIGDG DG	
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	DB 9; Length 239; 8e-115; 0; Indels 0; Gaps	Similarity 100.0%; Score 1274; DB 9; Length 239; Similarity 100.0%; Pred. No. 4.8e-115; 9; Conservative 0; Mismatches 0; Indels 0; Gaps WYSKGEELFTGWPILVELDGDWNGHKFSVSGEGEGDATYGKLTKFICTTGKLEVPWPT 60	Duery Match 100.0%; Score 1274; DB 9; Length 239; Best Local Similarity 100.0%; Pred. No. 4.8e-115; Matches 239; Conservative 0; Mismatches 0; Indels 0; Gaps 1 MYSKGEBLFTGVVPILVELDGDVNCHKFSVSGEGEDATYGKLTLKFICTTGKLPVPWPT 60	Duery Match 100.0%; Score 1274; DB 9; Length 239; Best Local Similarity 100.0%; Pred. No. 4.8e-115; Matches 239; Conservative 0; Mismatches 0; Indels 0; Gaps 1 MVSKGELFTGVVPILVELDGDVNGHKFSVSGEGEGDATYGKLIKFICTTGKLPVPWPT 60 IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII	Duery Match 100.0%; Score 1274; DB 9; Length 239; Best Local Similarity 100.0%; Pred. No. 4.8e-115; Matches 239; Conservative 0; Mismatches 0; Indels 0; Gaps 1 MVSKGEELFTGVVPILVELDGDVUGHFEVSGEGEGDATYGKITIKFICTTGKLPVPWPT 60

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FEATURE: OTHER INFORMATION: Aequorea victoria green fluorescent protein modified as des OTHER INFORMATION: in specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61 LVTTLSYGVQCFSRYPDHMKQHDFFKSAMPEGYVQERTIFFKDDGNYKTRAEVKFEGDTL 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     121 VNRIELKGIDFKEDGNILGHKLEYNYNSHNYYIMADKOKNGIKVNFKIRHNIEDGSVOLA 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            181 DHYQQNTPIGDGPVLLPDNHYLSTQSALSKDPNEKRDHMVLLGFVTAAGITLGMDELYK 239
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 2, Application US/09797496B
Publication No. US20030049597Al
GENERAL INFORMATION:
APPLICANT: Simon, Sanford M.
APPLICANT: Chen, Yu
TITLE OF INVENTION: Chimeric Fluorescent Enzymes and Uses Thereof CURRENT APPLICATION NUMBER: US/09/797,496B
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin version 3.1
SEQ ID NO 2.2
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                                                                                                                                                                                                       APPLICANT: REGENTS OF THE UNIVERSITY OF CALIFORNIA
APPLICANT: TSIEN, ROGER
APPLICANT: Campbell, Robert
APPLICANT: Campbell, Robert
TITLE OF INVENTION: NON-OLIGOMERIZING FLUORESCENT PROTEINS
FILE REFERENCE: REGEN1530-2
CURRENT APPLICATION NUMBER: US/09/866,538
CURRENT FILING DATE: 2001-05-24
NUMBER OF SEQ ID NOS: 29
SOFTWARE: PATENTIN VERSION 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 99.1%; Score 1263; DB 9;
Best Local Similarity 99.2%; Pred. No. 5.5e-114;
Matches 237; Conservative 1; Mismatches 1;
                                                                                                                                                     Sequence 4, Application US/09866538 Publication No. US20030032088A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT CRGANISM: Aequorea victoria US-09-866-538-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
Matches 237; Conserv
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US-09-866-538-4
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Patent No. US20020157120A1

Patent No. US20020157120A1

PAPLICANT: THE REGENTS OF THE UNIVERSITY OF CALIFORNIA

APPLICANT: Taien, Roger Y.

APPLICANT: Baird, Geoffrey

TITLE OF INVENTION: CIRCULARLY PERMUTED FLUORESCENT PROTEIN INDICATORS

FILE REFERENCE: REGEN1470-1

CURRENT APPLICATION NUMBER: US/09/999,745

CURRENT FILING DATE: 2001-10-23

PRIOR APPLICATION NUMBER: 09/316,920

PRIOR PILING DATE: 1999-05-21

NUMBER OF SEQ ID NOS: 67

SOFTWARE: PATENTIN Version 3.0

SEQ ID NO 4

LENGTH: 239
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Pred. No. 5.5e-114;
1; Mismatches 1;
                                                                                                                                                                                                                                                                                  Score 1266; DB 9;
Pred. No. 2.8e-114;
0; Mismatches 1;
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION:
FILE REFERENCE: 3759-0115P
CURRENT FILING DATE: 2001-06-19
NUMBER OF SEQ ID NOS: 24
SOFTWARE: Patentin version 3.0
LENGTHARE: 299
TYPE: PRT
                                                                                                                                                                                                                                                                                      Query Match 99.4%;
Best Local Similarity 99.6%;
Matches 238; Conservative
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Local Similarity 99.2%;
nes 237; Conservative
                                                                                                                                                                                                                              ; ORGANISM: Aequoria Victoria
US-09-887-784-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; ORGANISM: Aequorea victoria
US-09-999-745-4
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                                                                                                                                                                                        LVTTLSYGVQCFSRYPDHMKQHDFFKSAMPEGYVQERTIFFKDDGNYKTRAEVKFEGDTL 120
                                                                               LVTTLSYGVQCFSRYPDHMKQHDFFKSAMPEGYVQERTIFFKDDGNYKTRAEVKFEGDTL 120
                                                                                                                                                                                                                                           1 MVSKGEELFTGVVPILVELDGDVNGHKFSVSGEGGGDATYGKLTLKFICTTGKLPVPWPT 60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: TSLEE, ROGET
APPLICANT: TSLEE, ROBERT
TILLE OF INVENTION: MONOBERIC AND DIMERIC FLUORESCENT
TILLE OF INVENTION: MONOBERIC AND DIMERIC FLUORESCENT
TILLE OF INVENTION: MONOBERIC AND METHODS FOR MAKING SAME
TILLE OF INVENTION: DROTEIN VARIANTS AND METHODS FOR MAKING
FRILER PEFERENCE: UC683.1CP2CP1
CURRENT APPLICATION NUMBER: US/10/121,258
FRIOR FILLING DATE: 2001-02-04
PRIOR PLILING DATE: 2001-02-24
PRIOR FILLING DATE: 2001-05-24
NUMBER OF SEQ ID NOS: 78
SOFTWARE: FastSED for Windows Version 4.0
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US-10-121-258-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 7, Application US/10221461
Publication No. US20030092902A1
Eublication No. US20030092902A1
EUBLICANT: Marsh, Donald J.
TITLE OF INVENTION: MELANIN CONCENTRATING HORMONE RECEPTOR
TITLE OF INVENTION: CHIMBRIC AND FUSION PROTEINS
FILE REFERENCE: 20652P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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Pred. No. 5.5e-114;
1; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                   Sequence 13, Application US/10121258 Publication No. US20030059835A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 99.2
Matches 237; Conservative
                                                                                                                                                                                                                                                                                                                                                                      US-10-121-258-13
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61 LVTTLSYGVQCFSRYPDHMKQHDFFKSAMPEGYVQERTIFFKDDGNYKTRAEVKFEGDTL 120
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Squence 46, Application US/10100957A

Squence 46, Application US/10100957A

Publication No. US20030096322A1

GENERAL INFORMATION:

APPLICANT: Giullano, Kenneth A.

APPLICANT: Kapur, Ravi

TITLE OF INVENTION: A system for Cell Based Screening

FILE REFERENCE: 97-022-L1A

CURRENT FILIGE DATE: 2002-03-19

NUMBER OF SEQ ID NOS: 180

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 46

LENGTH: 239
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                                                                                                                                                                                                                                                                                                                                                                                      Query Match
99.1%; Score 1263; DB 9;
Best Local Similarity 99.2%; Pred. No. 5.5e-114;
Matches 237; Conservative 1; Mismatches 1:
CURRENT APPLICATION NUMBER: US/10/221,461
CURRENT FILING DATE: 2002-09-12
PRIOR APPLICATION NUMBER: PT/US01/08071
PRIOR FILING DATE: 2001-03-14
PRIOR PRIOR PELING DATE: 2000-03-15
NUMBER OF SEQ ID NOS: 37
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7.
                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: GFP derivative US-10-221-461-7
                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
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APPLICANT: BLAIR, DONALD
APPLICANT: CLAUSEN, PETER
APPLICANT: CLAUSEN, PETER
APPLICANT: CLAUSEN, PETER
APPLICANT: TOPOL, LILIA
APPLICANT: TOPOL, LILIA
APPLICANT: CALOTHY, GEORGES
TITLE OF INVENTION: WETHODS AND COMPOSITIONS FOR DRM, A SECRETED PROTEIN
TITLE OF INVENTION: WETHODS AND COMPOSITIONS FOR DRM, A SECRETED PROTEIN
TITLE OF INVENTION: WITH CELL GROWTH INHIBITING ACTIVITY
FILE REFERENCE: 14014.0356
CURRENT APPLICATION NUMBER: US/10/033,717
CURRENT FILING DATE: EARLIER FILING DATE: 1999-11-19
PRIOR PILING DATE: EARLIER FILING DATE: 1999-03-26
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/079,440
PRIOR FILING DATE: EARLIER FILING DATE: 1999-03-26
NUMBER OF SEQ ID MOS: 38
SOFTWARR: FASTSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ) OTHER INFORMATION: Description of Artificial Sequence:/No. US20030078406Ale is OTHER INFORMATION: synthetic construct US-10-033-717-35
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                                                                                                   121 VNRIELKGIDFKEDGNILGHKLEYNYNSHNYYIMADKQKNGIKVNFKIRHNIEDGSVQLA 180
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MASKGEELFTGVVPILVELDGDVNGHKFSVSGEGEGDATYGKLTLKFICTTGKLPVPWPT
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US-10-033-717-33
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LENGIH: 30
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PUblication No. US20030096322A1
GENERAL INFORMATION:
APPLICANT: GAlliano, Kenneth A.
APPLICANT: Kapur, Ravi
ITILE OF INVENTION: A System for Cell Based Screening
FILE REPERENCE: 97-022-LIA
CURRENT FILIAG DATE: 2002-03-19
NUMBER OF SEQ ID NOS: 180
SOFTWARE: Patentin Ver. 2.0
LENGTH: 294
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 99.1%; Score 1263; DB 10;
Best Local Similarity 99.2%; Pred. No. 5.5e-114;
Matches 237; Conservative 1; Mismatches 1;
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99.2%; Pred. No. 7.3e-114;
tive 1; Mismatches 1;
                                                                                                                                                                   GAPERAL INCORMATION:
APPLICANT: Miyawaki, Atsushi
APPLICANT: Sawano, Asako
TILLE OF INVENTION: METHOD FOR MUTACENESIS
FILE REFERENCE: 11283-012001
CURRENT APPLICATION WUMBER: US/09/920,922
CURRENT FILING DATE: 2001-08-02
PRIOR APPLICATION NUMBER: JP 2000-237166
PRIOR FILING DATE: 2000-08-04
PRIOR FILING DATE: 2000-08-04
SUMBER OF SEQ ID NOS: 9
SOFWMARE: PRACECO for Windows Version 4.0
                                                                                                                                     Sequence 2, Application US/09920922
Patent No. US20020083488A1
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; ORGANISM: Aequorea victoria
US-09-920-922-2
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Matches 237; Conservative
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US-10-100-957A-2
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LENGTH: 239
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US-10-033-717-34
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                                                         APPLICANT: BLAIR, DONALD
APPLICANT: CLAUSEN, PETER
APPLICANT: TOPOL, LILIA
APPLICANT: TOPOL, LILIA
APPLICANT: TOPOL, LILIA
APPLICANT: TOPOL, LILIA
APPLICANT: CALOTHY, GEORGES
TITLE OF INVENTION: WITH CELL GROWTH INHIBITING ACTIVITY
FILE REFERENCE: 14014.0356
CURRENT APPLICATION NUMBER: US/10/033,717
CURRENT APPLICATION NUMBER: 2001-12-27
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/444,066
PRIOR FILING DATE: EARLIER FILING DATE: 1999-03-11-19
PRIOR FILING DATE: EARLIER FILING DATE: 1999-03-26
PRIOR PELING DATE: EARLIER FILING DATE: 1998-03-26
PRIOR FILING DATE: EARLIER FILING DATE: 1998-03-26
NUMBER OF SEQ ID NOS: 38
SOFTWARE: FASTER APPLICATION NUMBER: 60/079,440
SPENDR FILING DATE: EARLIER FILING DATE: 1998-03-26
NUMBER OF SEQ ID NOS: 38
SEQ ID NO 33
SEQ ID NO 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: Description of Artificial Sequence:/No. US20030078406Ale; OTHER INFORMATION: synthetic construct
US-10-033-717-33
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APPLICANT: CLAUSEN, PETER
APPLICANT: TOPOL, LILIA
APPLICANT: MARTH MARTHODE
APPLICANT: CALOTHY, GEORGES
TITLE OF INVENTION: WETHODS AND COMPOSITIONS FOR DRM, A SECRETED PROTEIN
TITLE OF INVENTION: WITH CELL GROWTH INHIBITING ACTIVITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 MVSKGEELFTGVVPILVELDGDVNGHKFSVSGEGEGDATYGKLTLKFICTTGKLPVPWPT
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PRIOR FILING DATE: 2001-12-27
PRIOR FILING DATE: EARLIER APPLICATION NUMBER: 09/444,066
PRIOR FILING DATE: EARLIER FILING DATE: 1999-11-19
PRIOR PELICATION NUMBER: EARLIER APPLICATION NUMBER: 09/277,407
PRIOR PELICATION NUMBER: EARLIER APPLICATION NUMBER: 09/277,407
PRIOR PELICATION NUMBER: EARLIER FILING DATE: 1999-03-26
PRIOR FILING DATE: EARLIER FILING DATE: 1998-03-26
NUMBER OF SEQ ID NOS: 38
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Pred. No. 9.5e-114;
1; Mismatches 1;
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CURRENT APPLICATION NUMBER: US/10/033,717
CURRENT FILING DATE: 2001-12-27
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Publication No. US20030078406A1
GENERAL INFORMATION:
Application US/10033717 No. US20030078406A1
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99.2%;
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ORGANISM: Artificial Sequence
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237; Conservative
                                          GENERAL INFORMATION
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Best Local S
Matches 237
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    Sequence 33,
Publication
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TITLE OF INVENTION: A Method For Extracting Quantitative Information Relating TO TITLE OF INVENTION: On A Cellular Response FILE REFERENCE: 3759-0120P CURRENT APPLICATION NUMBER: US/10/072,036 CURRENT FILING DATE: 2002-09-13 PRIOR PAPILCATION NUMBER: 09/417,197 PRIOR FILING DATE: 1999-10-07 NUMBER: OF SEQ ID NOS: 143
                                                                                                             FEATURE: OTHER INFORMATION: Description of Artificial Sequence:/No. US20030078406Ale OTHER INFORMATION: synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                141 MySKGBELFTGVVPILVELDGDVNGHKFSVSGEGEGDATYGKLTLKFICTTGKLPVPWPT 200
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                                                                                                                                                                                                                                                                                                                                                                                                                                 VNRIELKGIDFKEDGNILGHKLEYNYNSHNYYIMADKOKNGIKVNFKIRHNIEDGSVOLA 180
                                                                                                                                                                                                                                                                                                                                                                                                       LVIILSYGVQCFSRYPDHMKQHDFFKSAMPEGYVQERIIFFKDDGNYKIRAEVKFEGDIL 120
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                                                                                                                                                                                                                         Length 359;
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                                                                                                                                                                                                                           Score 1263; DB 9;
Pred. No. 9.5e-114;
1; Mismatches 1;
SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 34 LENGTH: 359
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequence 129, Application US/10072036
publication No. US20030082564A1
general information:
APPLICANT: Ole THASTRUP
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Best Local Similarity 99.2%;
Matches 237; Conservative
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                                                                                                                                                                                                                           Query Match 99.1%;
Best Local Similarity 99.2%;
Matches 237; Conservative
                                                                       TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: SOTEN TULLIN
APPLICANT: Kasper ALMHOLT
APPLICANT: Kurt SCUDDER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Sara BJRON
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; OTHER INFORMATION: Description of Artificial Sequence:/No. US20030078406Ale = ; OTHER INFORMATION: synthetic construct US-10-033-717-32
                                                                                                                                                   Sequence 32, Application US/10033717

Sequence 32, Application No. US20030078406a1

GENERAL INFORMATION:
APPLICANT: CLAUSEN, PETER
APPLICANT: CLAUSEN, PETER
APPLICANT: CLAUSEN, PETER
APPLICANT: CALOTHY, GEORGES
TITLE OF INVENTION: WITH CELL GROWTH INHIBITING ACTIVITY
CURRENT APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/444,066
FRIOR PILING DATE: EARLIER FILING DATE: 1999-11-19
PRIOR FILING DATE: EARLIER FILING DATE: 1999-03-26
SPRIOR PELING DATE: EARLIER FILING DATE: 1999-03-26
NUMBER OF SEQ ID MOS: 38
SOFTWARE: FRABER FILING DATE: 1998-03-26
NUMBER OF SEQ ID MOS: 38
SECOFTWARE: FRABER FILING DATE: 1998-03-26
SED OF MOS: 38
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99.1%; Score 1263; DB 9; Length 3
Best Local Similarity 99.2%; Pred. No. 1.1e-113;
Matches 237; Conservative 1; Mismatches 1; Indels
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Search completed: June 3, 2003, 15:17:10 Job time : 46 secs